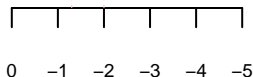
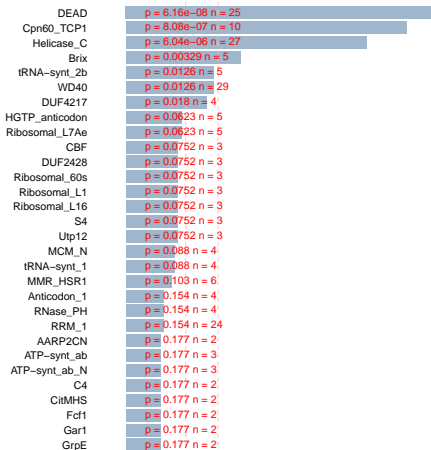


Bulk.Fem-Mal_difpos domain



log(p)

n = 1165 genes | p adjust BH TRUE